

Motion Estimation, Modeling and Compensation

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Introduction

MRI is intrinsically susceptible to motion artifacts; however, for the same reasons MRI offers numerous techniques for measuring motion¹. High performance MRI systems have increased the speed of data acquisition and thus reduced the susceptibility to unwanted motion artifacts, while increasing the accuracy of motion estimation. We are now at the limit of gradient switching speeds; peripheral nerve stimulation will not allow faster switching times unless some method is used to shield the patient from the stimulating field, so new techniques will have to work within this limitation.

CINE MRI Acquisition

One of the most effective methods for observing motion is the simplest: obtain a movie of the moving object. Even when the image quality of the individual frames is not great, the appreciation of the motion can be quite satisfactory. For example, CINE acquisitions of the heart can show significant wall motion abnormalities that are not appreciated from the individual still frames. Similarly, these movies of the heart can directly demonstrate asynchronous activation of the ventricle².

For a motion that is repeated, such as contraction of the heart or blood flow in major vessels, movies in multiple slices can be reconstructed from data obtained from 2-16 consecutive heartbeats during a patient breath-hold. If the patient can hold their diaphragm still for this long, and the ECG trigger works well for each beat, and each of the 2-16 beats is very similar, movies of extremely good quality are produced³⁻⁶. However, it is not uncommon for at least one of the above constraints to be violated in a sick patient. In this case, reduced data acquisition strategies such as TSENSE⁷ or UNFOLD^{8,9} can be employed to yield real-time movies¹⁰⁻¹².

Spin Tagging

The principle of "tagging" spins with a saturation pulse was first proposed by Morse and Singer to measure bulk flow¹³. It was demonstrated by Zerhouni et. al. that the same principle could be used to visually mark tissue with tagged magnetization to measure the more complex deformations of the heart¹⁴. Axel and Dougherty subsequently proposed a very efficient scheme for generating parallel planes of saturation throughout the entire imaging volume^{15,16}. Many investigators have proposed refinements and extensions of these methods for generating more complex saturation patterns¹⁷⁻²². Extensive efforts have been directed towards validating these patterns for measuring deformation with phantom experiments of both stationary^{23,24} and moving samples²⁵⁻²⁷.

We can break down the process of motion tracking with spin tagging into three stages: (1) a saturation pattern is placed in the myocardial tissue with spatially selective rf pulses, (2) a sequence of MR images is obtained in which the displacement of the saturation pattern can be observed, (3) the motion of the saturation pattern is used to solve for the motion of the tissue. Consideration of the rate and extent of the motion must be taken into account when designing the appropriate tagging pattern and imaging protocol²⁷. Because of the simple nature of the saturation pattern, virtually any imaging sequence can have tagging pulses added to it.

Imaging Tagged Magnetization

Although tagging pulses can be added to any imaging sequence, the imaging sequence must provide good contrast between the tagged and non-tagged tissue^{6;28} and provide the temporal and spatial resolution required to track the saturation patterns²⁷.

Probably the most common sequence used to image the tagging pattern is the segmented k-space GRE or SSFP cardiac gated CINE sequence. The advantage of segmented k-space imaging is that the signal is sampled within a brief readout window with very short TE, (utilizing fractional echoes). This reduces flow artifact and loss of signal due to magnetic susceptibility artifacts. The disadvantage of segmented k-space imaging is the relatively low efficiency of these methods because an rf pulse is required for each line of k-space²⁹. Also, the contrast between the tags and the background tissue decreases as both are driven towards the same steady-state with many imaging pulses^{6;28;30}. Echo planar imaging yields high efficiency, high tag contrast images; however, single shot EPI images of the heart are extremely difficult to obtain on a consistent basis because of local magnetic field inhomogeneities^{31;32}. A compromise of segmented echo-planar imaging^{33;34} allows us to tune the readout duration to its optimum. In the heart, the optimal total EPI readout duration is about 10ms. The addition of using TR values of less than 5 ms allows us to perform segmented EPI readouts while maintaining an SSFP type contrast and SNR^{5;6;35}. Through the variation of imaging tip angle, the user is able to tune the contrast to be more like bright blood SSFP (with lower tag contrast at later time points in the heart cycle), or darker blood SPGR (with extended tag contrast). The value of the optimal imaging tip angle will depend on the T1 and T2 of the tissue being imaged. If one were tracking displacement of the brain with SSFP tagging, the parameters would be significantly different than those in the heart.

Velocity Encoding Techniques

Measuring tissue motion with velocity encoding was originally proposed by Van Dijk³⁶ and further developed by Pelc et. al.³⁷ for CINE acquisitions. In contrast to the tagging methods which image prepared longitudinal magnetization, the velocity encoding methods obtain information about the dynamics of the tissue by phase encoding the velocity of the transverse magnetization shortly before the echo readout. This gives velocity encoding methods an advantage over tagging techniques because they do not suffer from tag pattern fading due to T1 recovery; however, they are more susceptible to motion artifacts. The development of fast switching gradients have reduced this susceptibility significantly, and remarkable datasets of blood flow patterns have been achieved³⁸. The trajectory of material points in the tissue of interest can be computed by integrating the three dimensional velocity field, or spatial gradients in the velocity field can be used to derive the strain-rate tensor in soft tissue directly^{37;39;40}.

Motion Tracking with Velocity Encoding Techniques

Because velocity encoding is a difference technique, at least four acquisitions are required to obtain the three dimensional velocity vector at each image pixel. Velocity encoding of the transverse magnetization is usually accomplished with bipolar flow encoding pulses. To obtain the three dimensional velocity at a specific point, the optimal combination of bipolar pulses is a tetrahedron⁴¹ but deviations from this pattern are possible depending on the circumstances¹². The different flow encoding acquisitions can be obtained in four contiguous TR periods, or obtained at the same time delay from the ECG in four heartbeats.

Harmonic Phase (HARP) Imaging Methods

In the HARP⁴² imaging method a sinusoidal tagging pattern is applied to the myocardium with a simple 1-1 SPAMM¹⁶ tagging pulse. This sinusoid moves the peak of the raw data to a region of k-space centered around the frequency of the sinusoid. If an image is reconstructed from a window of k-space around this peak, the magnitude of the image will be a low resolution picture of the amount of signal in the tagged sinusoid, and the *phase* of the image will show a banding pattern that deforms with the myocardial motion. Relative motion between movie frames of HARP images can be computed from the phase differences, hence motion can be measured without segmenting the position of the underlying tags. The elimination of the tag segmentation step in the analysis can reduce the analysis time by many orders of magnitude. The drawback is that the image is a relatively low resolution representation of the deformation field due to the fact that a window of k-space was used to reconstruct the HARP images. However, the fact that a small region of k-space is acquired means that the imaging can be performed very fast⁴³. The same encoding principle can be used to measure longitudinal strain while imaging in the short axis plane⁴⁴. This technique is particularly useful for measuring displacement fields that are spatially smooth.

Displacement Encoding using Stimulated Echoes (DENSE)

DENSE⁴⁵ imaging techniques use image phase to encode the net displacement of a pixel over time. Transverse magnetization is position-encoded by a short gradient pulse, stored as longitudinal magnetization during the “mixing time” (denoted TM), and then refocused before image data acquisition by an “unencoding” gradient pulse. The unencoding gradient pulse causes the transverse magnetization to refocus at a phase angle that is linearly proportional to the displacement that occurred during the mixing time. Unfortunately, only $\frac{1}{2}$ of the original transverse magnetization is recovered. However, the DENSE image has displacement estimates on the same spatial scale as the image pixel grid and those displacement estimates do not require image segmentation. These properties make DENSE a very attractive method for quantitative estimates of strain and displacement. Direct visualization of the myocardial deformation is not available, but it can be achieved by driving the motion of “synthetic” tags on the heart with the underlying measured displacement measurements. A number of variants of the DENSE technique have been developed to increase the SNR of the displacement estimates and reduce the effects of T1 recovery⁴⁶⁻⁵⁰. One drawback of DENSE is that it is a phase based technique. There are many causes of phase errors in imaging pixels around the heart, such as field

inhomogeneities, chemical shift, motion of the blood, and motion during breathing. Unfortunately, when the value of the phase of a pixel *is* the motion estimate, that estimate is susceptible to error from these many sources.

Modeling Motion

In order to obtain precise quantification of the tissue motion with spatial tagging techniques the position of the tags must be measured with a tag detection algorithm^{18;24;51;52}, and the three dimensional motions of the tissue computed from the combination of partial displacement information from each tag. A number of approaches can be employed to perform this estimation; most of the work to date has been done in modeling the motion of the heart⁵³⁻⁵⁶. General motion field reconstruction can be achieved with techniques such as B-splines in space and time⁵⁷⁻⁶⁰. Also, alternative methods exist for tracking motion such as optical flow⁶¹, HARP tagging⁴², and direct visualization of strain encoded MR⁴⁴. This motion modeling has been investigated extensively in modeling cardiac function⁶².

Motion Artifact Reduction

In many applications, patient motion is not the target, it is the problem. For example, high resolution imaging of the coronary arteries requires long acquisition times in order to achieve sufficient signal-to-noise within each small voxel. Unless we are able to increase the polarization of the coronary blood, data acquisition will likely remain in the “minutes” time frame for this application. Motion artifacts can appear due to displacement of the target from view to view; these can be reduced to varying degrees with phase encode view re-ordering methods⁶³⁻⁶⁵. For certain applications, the motion of the object from view to view can be determined from the imaging data itself⁶⁶⁻⁷⁰ or from specially crafted navigator pulses^{71;72}. Also, there is the potential for correcting image displacement in the spatial domain after full or partial image reconstruction^{73;74}.

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